



EBI Generic DB Entry Retrieval

ID AE007337 standard; DNA; PRO; 10216 BP.
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 AC AE007337; AE005672;
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 SV AE007337.1
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 DT 31-JUL-2001 (Rel. 68, Created)
 DT 02-SEP-2002 (Rel. 72, Last updated, Version 4)
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 DE Streptococcus pneumoniae TIGR4 section 20 of 194 of the complete genome.
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 KW .
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 OS Streptococcus pneumoniae TIGR4
 OC Bacteria; Firmicutes; Lactobacillales; Streptococcaceae; Streptococcus;
 OC Streptococcus pneumoniae.
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 RN [1]
 RP 1-10216
 RX MEDLINE; 21357209.
 RX PUBMED; 11463916.
 RA Tettelin H., Nelson K.E., Paulsen I.T., Eisen J.A., Read T.D., Peterson S.,
 RA Heidelberg J., DeBoy R.T., Haft D.H., Dodson R.J., Durkin A.S., Gwinn M.,
 RA Kolonay J.F., Nelson W.C., Peterson J.D., Umayam L.A., White O.,
 RA Salzberg S.L., Lewis M.R., Radune D., Holtzappple E., Khouri H., Wolf A.M.,
 RA Utterback T.R., Hansen C.L., McDonald L.A., Feldblyum T.V., Angiuoli S.,
 RA Dickinson T., Hickey E.K., Holt I.E., Loftus B.J., Yang F., Smith H.O.,
 RA Venter J.C., Dougherty B.A., Morrison D.A., Hollingshead S.K., Fraser C.M.;
 RT "Complete genome sequence of a virulent isolate of Streptococcus
 RT pneumoniae";
 RL Science 293(5529):498-506(2001).
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 RN [2]
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 RA Tettelin H., Nelson K.E., Paulsen I.T., Eisen J.A., Read T.D., Peterson S.,
 RA Heidelberg J., DeBoy R.T., Haft D.H., Dodson R.J., Durkin A.S., Gwinn M.,
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 RA Hickey E.K., Holt I.E., Loftus B.J., Ujwal M.L., Yang F., Smith H.O.,
 RA Venter J.C., Dougherty B.A., Morrison D.A., Hollingshead S.K., Fraser C.M.;
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 RL Submitted (29-JUN-2001) to the EMBL/GenBank/DDBJ databases.
 RL The Institute for Genomic Research, 9712 Medical Center Dr, Rockville, MD
 RL 20850, USA
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 DR SPTREMBL; Q97SS4; Q97SS4.
 DR SPTREMBL; Q97SS5; Q97SS5.
 DR SPTREMBL; Q97SS6; Q97SS6.

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General	Description	References	Links	Keywords	Sequence
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General information

Entry name	Q97ST2
Accession number	<u>Q97ST2</u>
Created	TrEMBLrel. 18, 1-OCT-2001
Sequence update	TrEMBLrel. 18, 1-OCT-2001
Annotation update	TrEMBLrel. 21, 1-JUN-2002

Description and origin of the Protein

Description	ABC transporter, ATP-binding protein.
Gene name(s)	SP0242.
Organism source	Streptococcus pneumoniae.
Taxonomy	Bacteria; Firmicutes; Bacillus/Clostridium group; Lactobacillales; Streptococcaceae; Streptococcus.
NCBI TaxID	<u>1313</u>

References

- [1] Tettelin,H., Nelson,K.E., Paulsen,I.T., Eisen,J.A., Read,T.D., Peterson,S., Heidelberg,J., DeBoy,R.T., Haft,D.H., Dodson,R.J., Durkin,A.S., Gwinn,M., Kolonay,J.F., Nelson,W.C., Peterson,J.D., Umayam,L.A., White,O., Salzberg,S.L., Lewis,M.R., Radune,D., Holtzapple,E., Khouri,H., Wolf,A.M., Utterback,T.R., Hansen,C.L., McDonald,L.A., Feldblyum,T.V., Angiuoli,S., Dickinson,T., Hickey,E.K., Holt,I.E., Loftus,B.J., Yang,F., Smith,H.O., Venter,J.C., Dougherty,B.A., Morrison,D.A., Hollingshead,S.K., Fraser,C.M.,
Complete genome sequence of a virulent isolate of Streptococcus pneumoniae.
(2001) *Science* **293**:498-506

Position	SEQUENCE FROM N.A.
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Comments	STRAIN=TIGR4;
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Medline	<u>21357209</u>
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PubMed	<u>11463916</u>
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Database cross-references

EMBL	<u>AE007337</u> ; <u>AAK74421.1</u> ; -.
TIGR	<u>SP0242</u> ; -.
InterPro	<u>IPR003593</u> ; AAA_ATPase. <u>IPR003439</u> ; ABC_transportr.
Pfam	<u>PF00005</u> ; ABC_tran; 1.
ProDom	<u>PD000006</u> ; ABC_transportr; 1.
SMART	<u>SM00382</u> ; AAA; 1.
PROSITE	<u>PS00211</u> ; ABC_TRANSPORTER; 1.

Keywords

ATP-binding; Complete proteome;**Sequence information****Length: 363 aa, molecular weight: 40757 Da, CRC64 checksum: 3D9F10B1D8DD7731**

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MQIAQYADRK	PDKLSGGQQQ	RVTLACALAV	NPSVLLMDEP	LSNLEAKLRL	DMRQAIREIQ	180
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HGT						363

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General	Description	References	Links	Keywords	Sequence
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